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## RAW SEQUENCE LISTING PATENT APPLICATION US/07/784,222

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This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING
2	(1)	eneral Information:
3 4	(1) G	eneral information.
5	(i)	APPLICANT: Westbrook, Carol A.
7 8 9	(ii)	TITLE OF INVENTION: Methods and Compositions for the Detection of Chromosomal Aberrations
10 11	(iii)	NUMBER OF SEQUENCES: 4
12 13 14 15 16 17 18	(iv)	CORRESPONDENCE ADDRESS:  (A) ADDRESSEE: Arnold, White & Durkee  (B) STREET: P. O. Box 4433  (C) CITY: Houston  (D) STATE: TX  (E) COUNTRY: USA  (F) ZIP: 77210-4433
20 21 22 23 24 25	(V)	COMPUTER READABLE FORM:  (A) MEDIUM TYPE: Floppy disk  (B) COMPUTER: IBM PC compatible  (C) OPERATING SYSTEM: PC-DOS/MS-DOS  (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
26 27 28 29 30	(vi)	CURRENT APPLICATION DATA: (A) APPLICATION NUMBER: US 07/784,222 (B) FILING DATE: 28-OCT-1991 (C) CLASSIFICATION:
31 32 33 34 35	(viii)	ATTORNEY/AGENT INFORMATION: (A) NAME: Parker, David L. (B) REGISTRATION NUMBER: 32,165 (C) REFERENCE/DOCKET NUMBER: ARCD:010\PAR
36 37 38 39	(ix)	TELECOMMUNICATION INFORMATION: (A) TELEPHONE: 713-787-1400 (B) TELEFAX: 713-789-2679
41 42	(2) INFO	RMATION FOR SEQ ID NO:1:
42 43 44 45 46	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single

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## RAW SEQUENCE LISTING PATENT APPLICATION US/07/784,222

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		INPU1 SE1: S1400.ra)
47	(D) TOPOLOGY: linear	
48 49	(ii) MOLECULE TYPE: DNA	
50	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
51 52	(XI) SEQUENCE DESCRIPTION. SEQ IS NO.1.	
53	AGGGTGCACA GCCGCAACGG C	21
54 55	(2) INFORMATION FOR SEQ ID NO:2:	
56	ALL STOURNER GUARAGERT CETGG.	
57 58	(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs	
59	(B) TYPE: nucleic acid	
60	(C) STRANDEDNESS: single	
61	(D) TOPOLOGY: linear	
62	(ii) VOI DOUTE MUDE. DNA	
63 64	(ii) MOLECULE TYPE: DNA	
65		
66	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
67		21
68	CGCTTCACTC AGACCCTGAG G	21
69 70		
71	(2) INFORMATION FOR SEQ ID NO:3:	
72	· ·	
73	(i) SEQUENCE CHARACTERISTICS:	
74 75	<ul><li>(A) LENGTH: 19 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	
76	(C) STRANDEDNESS: single	
77	(D) TOPOLOGY: linear	
78		
79	(ii) MOLECULE TYPE: DNA	
80 81		
82	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
83		19
84	GAAGGGCTTT TGAACTCTG	19
85 86		
87	(2) INFORMATION FOR SEQ ID NO:4:	
88	•	
89	(i) SEQUENCE CHARACTERISTICS:	
90	<ul><li>(A) LENGTH: 18 base pairs</li><li>(B) TYPE: nucleic acid</li></ul>	
91 92	(C) STRANDEDNESS: single	
93	(D) TOPOLOGY: linear	
94		
95	(ii) MOLECULE TYPE: DNA	
96 97	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
98	(VI) DEFORMED PROCETTITION. DRA IN NO. 4.	
99	GAAGCGCTTC TTCCTTAT	18

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## SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/07/784,222

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